

SEQUENCE LISTING

<110> Applied Research Systems ARS Holding N.V.

5 <120> NOVEL NOTCH-LIKE POLYPEPTIDES

<130> 809

10 <150> US60/436,785
<151> 2002-12-27

<160> 4

15 <170> PatentIn version 3.1

<210> 1
<211> 3711
<212> DNA
<213> homo sapiens

20 <400> 1
ctcacagcag cgcagcagag cgcagagcgg gctgccatgg cgctggccag gcctgggacc 60
ccggaccccc aggccctggc ctctgtcctg ctactgtctg tctgggcccc tgccctttcc 120
25 ctccctggctg ggacggtgcc ttcagagccc cccagtgcct gtgcctcaga cccgtgcgct 180
ccaggagaccg agtgccaggc taccgagagt ggtggctata cctgtgggac catggagccc 240
30 cgggggctgtg ccaccagacc atgccaccac ggcgctctgt gtgtgcccc a ggtccagat 300
cccaacggct tccgctgcta ctgcgtgccg ggtttccagg gccacgctg cgagctggac 360
atcgatgagt gtgcattccc gccgtgccac catggggcca cctgccgcaa cctggccgat 420
35 cgctacgagt gccattgccc ccttggttat gcaggcgtga cctgcgagat ggaggtggac 480
gagtgcgcct cagcgccctg cctgcacggg ggctcgtgcc tggacggcgt gggctccttc 540
40 cgctgtgtgt gcgcgccagg ctacgggggc acccgttgcc agctggacct cgacgagtgc 600
cagagccagc cgtgcgcaca tggggggcacg tgccacgacc tggtaaacgg gttccggtgc 660
gactgcgcgg gcaccggcta cgagggcacg cactgcgagc gggaggtgct ggagtgcgca 720
45 tcggcgccct gcgagcacia cgcgtcctgc ctgcaggggc tcgggagctt ccgctgcctc 780
tggtggccag gctacagcgg cgagctgtgc gaggtggacg aggacgagtg tgcattcgagc 840
50 ccctgccagc atggggggcc atgcctgcag cgctctgacc cggccctcta cgggggtgtc 900
caggccgcct tccctggcgc cttcagcttc cgccatgctg cgggtttcct gtgccactgc 960
ctcctggct ttgagggggc gacatgtgag gaagatgtgg atgaatgcct gtcggatccc 1020
55 tgcctgcacg gcggaacctg cagtgcacct gtggcaggct atatctgcag gtgccacagc 1080

	acctggggtg ggcgcgactg ttctgtgcag ctactggct gccagggcca cacctgcccg	1140
	ctggetgcca cctgcatccc tatcttcgag tctgggtcc acagttacgt ctgccactgc	1200
5	ccacctggta cccatggacc gttctgtggc cagaatacca cttctctgt gatggctggg	1260
	agccccattc aggcacagc gccagctggt ggccccctgg gtctggcact gaggtttcgc	1320
10	accacactgc ccgctgggac ctgggccact cgcaatgaca ccaaggaaag ctggagctg	1380
	gcattggtgg cagccacact tcaggccaca ctctggagct acagcaccac tgtgcttgc	1440
	ctgagactgc cggacctggc cctaaacgat ggccattggc accaggtgga ggttgtgctc	1500
15	catctagcga ccctggagct acggtctctg catgagggct gccctgcccg gctctgtgtg	1560
	gcctctggtc ctgtggccct ggcttccacg gcttcggcaa ctccgctgcc tgcggggtc	1620
20	tcctctgccc agctggggga cgcgacctt gcaggctgcc tccaggacgt gcgtgtggat	1680
	ggccacctcc tgctgcctga ggatctcgtg gagaacgtcc tcctgggctg tgagcgccga	1740
	gagcagtgcc ggctctgcc ttgtgtccac ggagggtcct gtgtggatct gtggactcat	1800
25	ttccgttgcg actgtgcccg gcccacataga ggtccacgt gcgtgatga gattcctgct	1860
	gccacctttg gcttgggagg cgcaccaagc tctgcctcct ttctgctcca agagctgcca	1920
30	ggtcccaacc tcacagtgc ttctcttctc cgcactcggg agtcgctgg cctgttgctc	1980
	cagtttgcca atgactccgc agctggccta acagtattcc tgagtgggg tcggatccgg	2040
	gctgaggcgc cgggcagtcc tgctgtagt ctccctgggc gctgggatga tgggctccgt	2100
35	cacctggtga tgctcagctt cgggcctgac cagctgcagg acctggggca gcacgtgcac	2160
	gtgggtggga ggctccttgc tgccgacagc cagccctggg gtgggcccct cagaggtgc	2220
40	ctccaggacc tgcgactcga tggctgccac ctcccctct ttctctgcc actggataac	2280
	tcaagccagc ccagcgagct cggcggcagg cagtccctga acctactgc gggctgcgtc	2340
	tccgaggaca tgtgcagtcc tgaccctgt ttcaatggtg ggacttgct cgtcacctgg	2400
45	aatgacttcc actgtacctg ccctgccaat ttacggggc ctacgtgtgc ccagcagctg	2460
	tggtgtcccc gccagccctg tctcccacct gccacgtgtg aggaggctcc tgatggcttt	2520
50	gtgtgtgtgg cggaggccac gtcccgcgag ggtcccccg ccgcgttcag cgggcacaac	2580
	gcgtcgtcag ggcgttgct cggcggtctg tcgtggcct ttgcacgcg cgactccgag	2640
	gcctggctgc tgctgcccgc ggcggggccc ctggaaggcg tgtggctggc ggtgcgcaat	2700
55	ggctcgtgg cggggggcgt gcggggaggc catggcctgc ccggcgctgt gctgcccata	2760
	ccggggccgc gcgtggccga tgggtgctgg caccgcgtgc gtctggccat ggagcgcccg	2820

gcggcgcgcca cctcgcgctg gctgctgtgg ctggatgggtg ccgccacccc ggtggcgctg 2880
 cgcggcctgg ccagtgaact gggcttcctg caggggcccg gtgctgtgcg catcctgctg 2940
 5 gctgagaact tcaccggctg cttggggccgc cacttgcct cttggcctgg gacgcgggcc 3000
 ccgatcctcg gctgccgggg cgcgcccggtg tgtgcgcct cgccctgtct gcacgacggt 3060
 10 gcctgccgtg acctcttcga cgcttttgcc tgcgcctgcg gccgggggtg ggaaggcccg 3120
 cgctgcgaag ccacgtcga ccctgtcac tccgccccct gcgcccggtg ccgctgtcac 3180
 acgcaccccg acggccgctt cgagtgcgcg tgcccgccctg gcttcggggg ccgcgcgtgc 3240
 15 aggttgctg tccatccaa ggagtgcagc ctgaatgtca cctgcctcga tggcagccca 3300
 tgtgaggggtg gctctcccgc tgccaactgc agctgcctgg agggctcttc tggccagagg 3360
 20 tgtcaggtcc cactctccc ctgtgaagcc aaccctgct tgaatggggg cacctgccgg 3420
 gcagctggag ggggtgtctga atgtatctgc aatgccagat tctccggcca gttctgtgaa 3480
 gtggcgaagg gcctgcccct gccgtgccca tcccactgc tggaggtggc cgtacctgca 3540
 25 gcctgtgctt gcctctcct cctcctcctg ggctctctt cagggatcct ggcagcccg 3600
 aagcgccgcc agtctgaggg cacctacagc ccaagccagc aggaggtggc tggggcccg 3660
 30 ctggagatgg acagtgtcct caaggtgccca ccggaggaga gactcatcta g 3711

<210> 2
 <211> 1236
 35 <212> PRT
 <213> homo sapiens
 <400> 2

40 Leu Thr Ala Ala Glu Gln Ser Ala Glu Arg Ala Ala Met Ala Leu Ala
 1 5 10 15
 45 Arg Pro Gly Thr Pro Asp Pro Gln Ala Leu Ala Ser Val Leu Leu Leu
 20 25 30
 50 Leu Leu Trp Ala Pro Ala Leu Ser Leu Leu Ala Gly Thr Val Pro Ser
 35 40 45
 Glu Pro Pro Ser Ala Cys Ala Ser Asp Pro Cys Ala Pro Gly Thr Glu
 50 55 60
 55 Cys Gln Ala Thr Glu Ser Gly Gly Tyr Thr Cys Gly Pro Met Glu Pro
 65 70 75 80

Arg Gly Cys Ala Thr Gln Pro Cys His His Gly Ala Leu Cys Val Pro
 85 90 95

5 Gln Gly Pro Asp Pro Asn Gly Phe Arg Cys Tyr Cys Val Pro Gly Phe
 100 105 110

10 Gln Gly Pro Arg Cys Glu Leu Asp Ile Asp Glu Cys Ala Ser Arg Pro
 115 120 125

15 Cys His His Gly Ala Thr Cys Arg Asn Leu Ala Asp Arg Tyr Glu Cys
 130 135 140

20 His Cys Pro Leu Gly Tyr Ala Gly Val Thr Cys Glu Met Glu Val Asp
 145 150 155 160

Glu Cys Ala Ser Ala Pro Cys Leu His Gly Gly Ser Cys Leu Asp Gly
 165 170 175

25 Val Gly Ser Phe Arg Cys Val Cys Ala Pro Gly Tyr Gly Gly Thr Arg
 180 185 190

30 Cys Gln Leu Asp Leu Asp Glu Cys Gln Ser Gln Pro Cys Ala His Gly
 195 200 205

35 Gly Thr Cys His Asp Leu Val Asn Gly Phe Arg Cys Asp Cys Ala Gly
 210 215 220

40 Thr Gly Tyr Glu Gly Thr His Cys Glu Arg Glu Val Leu Glu Cys Ala
 225 230 235 240

Ser Ala Pro Cys Glu His Asn Ala Ser Cys Leu Glu Gly Leu Gly Ser
 245 250 255

45 Phe Arg Cys Leu Cys Trp Pro Gly Tyr Ser Gly Glu Leu Cys Glu Val
 260 265 270

50 Asp Glu Asp Glu Cys Ala Ser Ser Pro Cys Gln His Gly Gly Arg Cys
 275 280 285

55 Leu Gln Arg Ser Asp Pro Ala Leu Tyr Gly Gly Val Gln Ala Ala Phe
 290 295 300

Pro Gly Ala Phe Ser Phe Arg His Ala Ala Gly Phe Leu Cys His Cys

	305		310		315		320									
5	Pro	Pro	Gly	Phe	Glu	Gly	Pro	Thr	Cys	Glu	Glu	Asp	Val	Asp	Glu	Cys
					325					330					335	
10	Leu	Ser	Asp	Pro	Cys	Leu	His	Gly	Gly	Thr	Cys	Ser	Asp	Thr	Val	Ala
				340					345					350		
15	Gly	Tyr	Ile	Cys	Arg	Cys	Pro	Glu	Thr	Trp	Gly	Gly	Arg	Asp	Cys	Ser
			355					360					365			
20	Val	Gln	Leu	Thr	Gly	Cys	Gln	Gly	His	Thr	Cys	Pro	Leu	Ala	Ala	Thr
		370					375					380				
25	Cys	Ile	Pro	Ile	Phe	Glu	Ser	Gly	Val	His	Ser	Tyr	Val	Cys	His	Cys
	385					390					395				400	
30	Pro	Pro	Gly	Thr	His	Gly	Pro	Phe	Cys	Gly	Gln	Asn	Thr	Thr	Phe	Ser
					405					410					415	
35	Val	Met	Ala	Gly	Ser	Pro	Ile	Gln	Ala	Ser	Val	Pro	Ala	Gly	Gly	Pro
			420					425						430		
40	Leu	Gly	Leu	Ala	Leu	Arg	Phe	Arg	Thr	Thr	Leu	Pro	Ala	Gly	Thr	Leu
		435						440					445			
45	Ala	Thr	Arg	Asn	Asp	Thr	Lys	Glu	Ser	Leu	Glu	Leu	Ala	Leu	Val	Ala
		450					455					460				
50	Ala	Thr	Leu	Gln	Ala	Thr	Leu	Trp	Ser	Tyr	Ser	Thr	Thr	Val	Leu	Val
		465				470					475				480	
55	Leu	Arg	Leu	Pro	Asp	Leu	Ala	Leu	Asn	Asp	Gly	His	Trp	His	Gln	Val
					485				490					495		
60	Glu	Val	Val	Leu	His	Leu	Ala	Thr	Leu	Glu	Leu	Arg	Leu	Trp	His	Glu
			500						505					510		
65	Gly	Cys	Pro	Ala	Arg	Leu	Cys	Val	Ala	Ser	Gly	Pro	Val	Ala	Leu	Ala
		515						520					525			
70	Ser	Thr	Ala	Ser	Ala	Thr	Pro	Leu	Pro	Ala	Gly	Ile	Ser	Ser	Ala	Gln
		530					535				540					

Leu Gly Asp Ala Thr Phe Ala Gly Cys Leu Gln Asp Val Arg Val Asp
 545 550 555 560
 5 Gly His Leu Leu Leu Pro Glu Asp Leu Gly Glu Asn Val Leu Leu Gly
 565 570 575
 10 Cys Glu Arg Arg Glu Gln Cys Arg Pro Leu Pro Cys Val His Gly Gly
 580 585 590
 15 Ser Cys Val Asp Leu Trp Thr His Phe Arg Cys Asp Cys Ala Arg Pro
 595 600 605
 His Arg Gly Pro Thr Cys Ala Asp Glu Ile Pro Ala Ala Thr Phe Gly
 610 615 620
 20 Leu Gly Gly Ala Pro Ser Ser Ala Ser Phe Leu Leu Gln Glu Leu Pro
 625 630 635 640
 25 Gly Pro Asn Leu Thr Val Ser Phe Leu Leu Arg Thr Arg Glu Ser Ala
 645 650 655
 30 Gly Leu Leu Leu Gln Phe Ala Asn Asp Ser Ala Ala Gly Leu Thr Val
 660 665 670
 35 Phe Leu Ser Glu Gly Arg Ile Arg Ala Glu Ala Pro Gly Ser Pro Ala
 675 680 685
 Val Val Leu Pro Gly Arg Trp Asp Asp Gly Leu Arg His Leu Val Met
 690 695 700
 40 Leu Ser Phe Gly Pro Asp Gln Leu Gln Asp Leu Gly Gln His Val His
 705 710 715 720
 45 Val Gly Gly Arg Leu Leu Ala Ala Asp Ser Gln Pro Trp Gly Gly Pro
 725 730 735
 50 Phe Arg Gly Cys Leu Gln Asp Leu Arg Leu Asp Gly Cys His Leu Pro
 740 745 750
 55 Phe Phe Pro Leu Pro Leu Asp Asn Ser Ser Gln Pro Ser Glu Leu Gly
 755 760 765
 Gly Arg Gln Ser Trp Asn Leu Thr Ala Gly Cys Val Ser Glu Asp Met

	770	775	780
5	Cys Ser Pro Asp Pro Cys Phe Asn Gly Gly Thr Cys Leu Val Thr Trp 785 790 795 800		
10	Asn Asp Phe His Cys Thr Cys Pro Ala Asn Phe Thr Gly Pro Thr Cys 805 810 815		
15	Ala Gln Gln Leu Trp Cys Pro Gly Gln Pro Cys Leu Pro Pro Ala Thr 820 825 830		
20	Cys Glu Glu Val Pro Asp Gly Phe Val Cys Val Ala Glu Ala Thr Phe 835 840 845		
25	Arg Glu Gly Pro Pro Ala Ala Phe Ser Gly His Asn Ala Ser Ser Gly 850 855 860		
30	Arg Leu Leu Gly Gly Leu Ser Leu Ala Phe Arg Thr Arg Asp Ser Glu 865 870 875 880		
35	Ala Trp Leu Leu Arg Ala Ala Ala Gly Ala Leu Glu Gly Val Trp Leu 885 890 895		
40	Ala Val Arg Asn Gly Ser Leu Ala Gly Gly Val Arg Gly Gly His Gly 900 905 910		
45	Leu Pro Gly Ala Val Leu Pro Ile Pro Gly Pro Arg Val Ala Asp Gly 915 920 925		
50	Ala Trp His Arg Val Arg Leu Ala Met Glu Arg Pro Ala Ala Ala Thr 930 935 940		
55	Ser Arg Trp Leu Leu Trp Leu Asp Gly Ala Ala Thr Pro Val Ala Leu 945 950 955 960		
	Arg Gly Leu Ala Ser Asp Leu Gly Phe Leu Gln Gly Pro Gly Ala Val 965 970 975		
	Arg Ile Leu Leu Ala Glu Asn Phe Thr Gly Cys Leu Gly Arg His Phe 980 985 990		
	Ala Ser Trp Pro Gly Thr Pro Ala Pro Ile Leu Gly Cys Arg Gly Ala 995 1000 1005		

Pro Val Cys Ala Pro Ser Pro Cys Leu His Asp Gly Ala Cys Arg
 1010 1015 1020
 5 Asp Leu Phe Asp Ala Phe Ala Cys Ala Cys Gly Pro Gly Trp Glu
 1025 1030 1035
 10 Gly Pro Arg Cys Glu Ala His Val Asp Pro Cys His Ser Ala Pro
 1040 1045 1050
 15 Cys Ala Arg Gly Arg Cys His Thr His Pro Asp Gly Arg Phe Glu
 1055 1060 1065
 20 Cys Arg Cys Pro Pro Gly Phe Gly Gly Pro Arg Cys Arg Leu Pro
 1070 1075 1080
 Val Pro Ser Lys Glu Cys Ser Leu Asn Val Thr Cys Leu Asp Gly
 1085 1090 1095
 25 Ser Pro Cys Glu Gly Gly Ser Pro Ala Ala Asn Cys Sér Cys Leu
 1100 1105 1110
 30 Glu Gly Leu Ala Gly Gln Arg Cys Gln Val Pro Thr Leu Pro Cys
 1115 1120 1125
 35 Glu Ala Asn Pro Cys Leu Asn Gly Gly Thr Cys Arg Ala Ala Gly
 1130 1135 1140
 40 Gly Val Ser Glu Cys Ile Cys Asn Ala Arg Phe Ser Gly Gln Phe
 1145 1150 1155
 Cys Glu Val Ala Lys Gly Leu Pro Leu Pro Leu Pro Phe Pro Leu
 1160 1165 1170
 45 Leu Glu Val Ala Val Pro Ala Ala Cys Ala Cys Leu Leu Leu Leu
 1175 1180 1185
 50 Leu Leu Gly Leu Leu Ser Gly Ile Leu Ala Ala Arg Lys Arg Arg
 1190 1195 1200
 55 Gln Ser Glu Gly Thr Tyr Ser Pro Ser Gln Gln Glu Val Ala Gly
 1205 1210 1215
 Ala Arg Leu Glu Met Asp Ser Val Leu Lys Val Pro Pro Glu Glu

	1220	1225	1230	
5	Arg Leu Ile			
	1235			
	<210> 3			
	<211> 3570			
10	<212> DNA			
	<213> homo sapiens			
	<400> 3			
15	tcagagcccc ccagtgcctg tgcctcagac ccgtgcgctc cagggaccga gtgccaggct		60	
	accgagagtg gtggctatac ctgtgggccc atggagcccc ggggctgtgc caccagcca		120	
	tgccaccacg gcgctctgtg tgtgccccag ggtccagatc ccaacggctt ccgtgctac		180	
20	tgcgtgccgg gtttccaggg cccacgctgc gagctggaca tcgatgagtg tgcattccgg		240	
	ccgtgccacc atggggccac ctgccgaac ctggccgata gctacgagtg ccattgcccc		300	
25	cttggctatg caggcgtgac ctgcgagatg gaggtggacg agtgcgcctc agcgccctgc		360	
	ctgcacgggg gctcgtgcct ggacggcggt ggctccttcc gctgtgtgtg cgcgccaggc		420	
	tacgggggca cccgttgcca gctggacctc gacgagtgcc agagccagcc gtgcgcacat		480	
30	gggggcacgt gccacgacct ggtcaacggg ttccgggtcg actgcgcggg caccggctac		540	
	gagggcacgc actgcgagcg ggaggtgctg gactgcgcat cggcgccctg cgagcacaac		600	
	gcgtcctgcc tcgagggcct cgggagcttc cgctgcctct gttggccagg ctacagcggc		660	
35	gagctgtgcg aggtggacga ggacgagtgt gcacgagcc cctgccagca tgggggcca		720	
	tgcctgcagc gctctgaccc ggccctctac gggggtgtcc aggccgcctt ccctggcgcc		780	
40	ttcagcttcc gccatgctgc gggtttcctg tgccactgcc ctcttggtt tgaggggccg		840	
	acatgtgagg aagatgtgga tgaatgcctg tcggatccct gcctgcacgg cggaacctgc		900	
	agtgcactg tggcaggcta tatctgcagg tgcccagaga cctgggggtg gcgcgactgt		960	
45	tctgtgcagc tcaactggctg ccagggccac acctgcccgc tggetgccac ctgcatccct		1020	
	atcttcgagt ctgggggtcca cagttacgtc tgccactgcc cactgggtac ccatggaccg		1080	
50	ttctgtggcc agaataccac cttctctgtg atggctggga gcccattca ggcatcagt		1140	
	ccagctgggt gcccctggg tctggcactg aggtttcgca ccacactgcc cgctgggacc		1200	
	ttggccactc gcaatgacac caaggaaagc ttggagctgg cattggtggc agccacactt		1260	
55	caggccacac tctggagcta cagcaccact gtgctgtgcc tgagactgcc ggacctggcc		1320	
	ctaaacgatg gccattggca ccaggtggag gttgtgtctc atctagcgac cctggagcta		1380	

	cggtctctggc atgagggctg ccctgcccgg ctctgtgtgg cctctggtcc tgtggccctg	1440
5	gcttccacgg cttcggcaac tccgctgcct gccgggatct cctctgcca gctgggggac	1500
	gcgacctttg caggctgcct ccaggacgtg cgtgtggatg gccacctcct gctgcctgag	1560
	gatctcggtg agaacgtcct cctgggctgt gagcgccgag agcagtgcg gcctctgcct	1620
10	tgtgtccacg gagggctcctg tgtggatctg tggactcatt tccgttgcca ctgtgcccg	1680
	ccccatagag gtcccacgtg cgtgatgag attcctgctg ccacctttgg cttgggaggc	1740
	gccccaaagt ctgcctcctt tctgctccaa gagctgccag gtcccaacct cacagtgtct	1800
15	ttccttctcc gcaactcggga gtccgctggc ctgttgctcc agtttgccaa tgactccgca	1860
	gctggcctaa cagtattcct gagtgggggt cggatccggg ctgaggcgcc gggcagtcct	1920
20	gctgtagtgc tccctgggag ctgggatgat gggctccgtc acctggtgat gctcagcttc	1980
	gggcctgacc agctgcagga cctggggcag cacgtgcacg tgggtgggag gctccttgct	2040
25	gccgacagcc agccctgggg tgggcccctc cgaggctgcc tccaggacct gcgactcgat	2100
	ggctgccacc tccccttctt tcctctgcca ctggataact caagccagcc cagcgagctc	2160
	ggcggcaggc agtcctggaa cctcactgag ggctgcgtct ccgaggacat gtgcagtcct	2220
30	gacccctgtt tcaatggtgg gacttgcttc gtcacctgga atgacttcca ctgtacctgc	2280
	cctgccaatt tcacggggcc tacgtgtgcc cagcagctgt ggtgtcccgg ccagccctgt	2340
35	ctcccacctg ccacgtgtga ggaggctcct gatggctttg tgtgtgtggc ggaggccacg	2400
	ttccgcgagg gtccccccgc cgcgttcagc gggcacaacg cgtcgtcagg gcgcttgctc	2460
	ggcggcctgt cgtggcctt tcgcacgcgc gactccgagg cctggctgct gcgtgccgcg	2520
40	gcgggcgccc tggaaggcgt gtggctggcg gtgcgcaatg gctcgtggc ggggggcgtg	2580
	cgcggaggcc atggcctgcc cggcgtgtg ctgcccatac cggggccgcg cgtggccgat	2640
45	ggtgcctggc accgcgtgcg tctggccatg gagcgcccgg cggccgccac ctgcgctgg	2700
	ctgctgtggc tggatggtgc cgccaccccg gtggcgtgc gcggcctggc cagtgcctg	2760
	ggcttctgc agggcccggtg tgctgtgcgc atcctgctgg ctgagaactt caccggtgc	2820
50	ttgggcgcgc acttcgcctc ttggcctggg acgcccggcc cgatcctcgg ctgccgcggc	2880
	gcgcccgtgt gtgcgcctc gccctgtctg cacgacggtg cctgccgtga cctcttcgac	2940
55	gcctttgcct gcgcctgcg cccgggggtg gaaggcccgc gctgcgaagc ccacgtcgac	3000
	ccctgtcact ccgcccctg cgcctgtggc cgtgtgcaca cgcacccga cggccgcttc	3060
	gagtgcgct gcccgccctg cttcgggggc ccgcgctgca ggttgccctgt cccatccaag	3120

gagtgcagcc tgaatgtcac ctgcctcgat ggcagcccat gtgaggggtgg ctctcccgt 3180
 gccaaactgca gctgcctgga gggctcttgcgt ggccagaggt gtcagggtccc cactctcccc 3240
 5 tgtgaagcca acccctgctt gaatgggggc acctgccggg cagctggagg ggtgtctgaa 3300
 tgtatctgca atgccagatt ctccggccag ttctgtgaag tggcgaaggg cctgccccctg 3360
 10 ccgctgccat tcccactgct ggaggtggcc gtacctgcag cctgtgcctg cctcctcctc 3420
 ctctcctctgg gcctcctttc agggatcctg gcagcccgaa agcgccgcca gtctgagggc 3480
 acctacagcc caagccagca ggaggtggct ggggcccggc tggagatgga cagtgtcctc 3540
 15 aaggtgccac cggaggagag actcatctag 3570

<210> 4
 20 <211> 1189
 <212> PRT
 <213> homo sapiens

<400> 4
 25 Ser Glu Pro Pro Ser Ala Cys Ala Ser Asp Pro Cys Ala Pro Gly Thr
 1 5 10 15

30 Glu Cys Gln Ala Thr Glu Ser Gly Gly Tyr Thr Cys Gly Pro Met Glu
 20 25 30

35 Pro Arg Gly Cys Ala Thr Gln Pro Cys His His Gly Ala Leu Cys Val
 35 40 45

40 Pro Gln Gly Pro Asp Pro Asn Gly Phe Arg Cys Tyr Cys Val Pro Gly
 50 55 60

45 Phe Gln Gly Pro Arg Cys Glu Leu Asp Ile Asp Glu Cys Ala Ser Arg
 65 70 75 80

50 Pro Cys His His Gly Ala Thr Cys Arg Asn Leu Ala Asp Arg Tyr Glu
 85 90 95

55 Cys His Cys Pro Leu Gly Tyr Ala Gly Val Thr Cys Glu Met Glu Val
 100 105 110

Asp Glu Cys Ala Ser Ala Pro Cys Leu His Gly Gly Ser Cys Leu Asp
 115 120 125

Gly Val Gly Ser Phe Arg Cys Val Cys Ala Pro Gly Tyr Gly Gly Thr

	130	135	140
5	Arg Cys Gln Leu Asp Leu Asp Glu Cys Gln Ser Gln Pro Cys Ala His 145 150 155 160		
10	Gly Gly Thr Cys His Asp Leu Val Asn Gly Phe Arg Cys Asp Cys Ala 165 170 175		
15	Gly Thr Gly Tyr Glu Gly Thr His Cys Glu Arg Glu Val Leu Glu Cys 180 185 190		
20	Ala Ser Ala Pro Cys Glu His Asn Ala Ser Cys Leu Glu Gly Leu Gly 195 200 205		
25	Ser Phe Arg Cys Leu Cys Trp Pro Gly Tyr Ser Gly Glu Leu Cys Glu 210 215 220		
30	Val Asp Glu Asp Glu Cys Ala Ser Ser Pro Cys Gln His Gly Gly Arg 225 230 235 240		
35	Cys Leu Gln Arg Ser Asp Pro Ala Leu Tyr Gly Gly Val Gln Ala Ala 245 250 255		
40	Phe Pro Gly Ala Phe Ser Phe Arg His Ala Ala Gly Phe Leu Cys His 260 265 270		
45	Cys Pro Pro Gly Phe Glu Gly Pro Thr Cys Glu Glu Asp Val Asp Glu 275 280 285		
50	Cys Leu Ser Asp Pro Cys Leu His Gly Gly Thr Cys Ser Asp Thr Val 290 295 300		
55	Ala Gly Tyr Ile Cys Arg Cys Pro Glu Thr Trp Gly Gly Arg Asp Cys 305 310 315 320		
	Ser Val Gln Leu Thr Gly Cys Gln Gly His Thr Cys Pro Leu Ala Ala 325 330 335		
	Thr Cys Ile Pro Ile Phe Glu Ser Gly Val His Ser Tyr Val Cys His 340 345 350		
	Cys Pro Pro Gly Thr His Gly Pro Phe Cys Gly Gln Asn Thr Thr Phe 355 360 365		

Ser Val Met Ala Gly Ser Pro Ile Gln Ala Ser Val Pro Ala Gly Gly
 370 375 380

5
 Pro Leu Gly Leu Ala Leu Arg Phe Arg Thr Thr Leu Pro Ala Gly Thr
 385 390 395 400

10
 Leu Ala Thr Arg Asn Asp Thr Lys Glu Ser Leu Glu Leu Ala Leu Val
 405 410 415

15
 Ala Ala Thr Leu Gln Ala Thr Leu Trp Ser Tyr Ser Thr Thr Val Leu
 420 425 430

20
 Val Leu Arg Leu Pro Asp Leu Ala Leu Asn Asp Gly His Trp His Gln
 435 440 445

25
 Val Glu Val Val Leu His Leu Ala Thr Leu Glu Leu Arg Leu Trp His
 450 455 460

30
 Glu Gly Cys Pro Ala Arg Leu Cys Val Ala Ser Gly Pro Val Ala Leu
 465 470 475 480

35
 Ala Ser Thr Ala Ser Ala Thr Pro Leu Pro Ala Gly Ile Ser Ser Ala
 485 490 495

40
 Gln Leu Gly Asp Ala Thr Phe Ala Gly Cys Leu Gln Asp Val Arg Val
 500 505 510

45
 Asp Gly His Leu Leu Leu Pro Glu Asp Leu Gly Glu Asn Val Leu Leu
 515 520 525

50
 Gly Cys Glu Arg Arg Glu Gln Cys Arg Pro Leu Pro Cys Val His Gly
 530 535 540

55
 Gly Ser Cys Val Asp Leu Trp Thr His Phe Arg Cys Asp Cys Ala Arg
 545 550 555 560

60
 Pro His Arg Gly Pro Thr Cys Ala Asp Glu Ile Pro Ala Ala Thr Phe
 565 570 575

65
 Gly Leu Gly Gly Ala Pro Ser Ser Ala Ser Phe Leu Leu Gln Glu Leu
 580 585 590

Pro Gly Pro Asn Leu Thr Val Ser Phe Leu Leu Arg Thr Arg Glu Ser
 595 600 605
 5 Ala Gly Leu Leu Leu Gln Phe Ala Asn Asp Ser Ala Ala Gly Leu Thr
 610 615 620
 10 Val Phe Leu Ser Glu Gly Arg Ile Arg Ala Glu Ala Pro Gly Ser Pro
 625 630 635 640
 15 Ala Val Val Leu Pro Gly Arg Trp Asp Asp Gly Leu Arg His Leu Val
 645 650 655
 Met Leu Ser Phe Gly Pro Asp Gln Leu Gln Asp Leu Gly Gln His Val
 660 665 670
 20 His Val Gly Gly Arg Leu Leu Ala Ala Asp Ser Gln Pro Trp Gly Gly
 675 680 685
 25 Pro Phe Arg Gly Cys Leu Gln Asp Leu Arg Leu Asp Gly Cys His Leu
 690 695 700
 30 Pro Phe Phe Pro Leu Pro Leu Asp Asn Ser Ser Gln Pro Ser Glu Leu
 705 710 715 720
 35 Gly Gly Arg Gln Ser Trp Asn Leu Thr Ala Gly Cys Val Ser Glu Asp
 725 730 735
 Met Cys Ser Pro Asp Pro Cys Phe Asn Gly Gly Thr Cys Leu Val Thr
 740 745 750
 40 Trp Asn Asp Phe His Cys Thr Cys Pro Ala Asn Phe Thr Gly Pro Thr
 755 760 765
 45 Cys Ala Gln Gln Leu Trp Cys Pro Gly Gln Pro Cys Leu Pro Pro Ala
 770 775 780
 50 Thr Cys Glu Glu Val Pro Asp Gly Phe Val Cys Val Ala Glu Ala Thr
 785 790 795 800
 55 Phe Arg Glu Gly Pro Pro Ala Ala Phe Ser Gly His Asn Ala Ser Ser
 805 810 815
 Gly Arg Leu Leu Gly Gly Leu Ser Leu Ala Phe Arg Thr Arg Asp Ser
 820 825 830

5 Glu Ala Trp Leu Leu Arg Ala Ala Ala Gly Ala Leu Glu Gly Val Trp
 835 840 845

 10 Leu Ala Val Arg Asn Gly Ser Leu Ala Gly Gly Val Arg Gly Gly His
 850 855 860

 15 Gly Leu Pro Gly Ala Val Leu Pro Ile Pro Gly Pro Arg Val Ala Asp
 865 870 875 880

 20 Gly Ala Trp His Arg Val Arg Leu Ala Met Glu Arg Pro Ala Ala Ala
 885 890 895

 25 Thr Ser Arg Trp Leu Leu Trp Leu Asp Gly Ala Ala Thr Pro Val Ala
 900 905 910

 30 Leu Arg Gly Leu Ala Ser Asp Leu Gly Phe Leu Gln Gly Pro Gly Ala
 915 920 925

 35 Val Arg Ile Leu Leu Ala Glu Asn Phe Thr Gly Cys Leu Gly Arg His
 930 935 940

 40 Phe Ala Ser Trp Pro Gly Thr Pro Ala Pro Ile Leu Gly Cys Arg Gly
 945 950 955 960

 45 Ala Pro Val Cys Ala Pro Ser Pro Cys Leu His Asp Gly Ala Cys Arg
 965 970 975

 50 Asp Leu Phe Asp Ala Phe Ala Cys Ala Cys Gly Pro Gly Trp Glu Gly
 980 985 990

 55 Pro Arg Cys Glu Ala His Val Asp Pro Cys His Ser Ala Pro Cys Ala
 995 1000 1005

 60 Arg Gly Arg Cys His Thr His Pro Asp Gly Arg Phe Glu Cys Arg
 1010 1015 1020

 65 Cys Pro Pro Gly Phe Gly Gly Pro Arg Cys Arg Leu Pro Val Pro
 1025 1030 1035

 70 Ser Lys Glu Cys Ser Leu Asn Val Thr Cys Leu Asp Gly Ser Pro
 1040 1045 1050

